DT05 Rec'd PCT/PT0 1 0 DEC 2004

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125

Asp Cys Leu Ile Tyr Lys Val Lys Val Leu Gly Gly Asn Phe Pro Ala

120

115

Asn	Gly	Pro	Val	Met	Gln	Lys	Lys	Ser	Lys	Gly	Trp	Glu	Pro	Cys	Thr	
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Glu	Met	Leu	Tyr	Pro	Arg	Asp	Gly	Val	Leu	Cys	Gly	Gln	Thr	Leu	Met	
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Ala	Leu	Lys	Cys	Ala	Asn	Gly	Lys	Asn	Leu	Thr	Cys	Gln	Leu	Arg	Ser	
				165					170					175		
Thr	Tyr	Arg	Ser	Arg	Lys	Gln	Ala	Ser	Ala	Leu	Gln	Thr	Pro	Gly	Phe	
			180					185					190			
His	Phe	Gly	Asp	His	Arg	Ile	Glu	Ile	Leu	Lys	Glu	Ala	Glu	Gly	Gly	
		195					200					205				
Asn	Tyr	Phe	Glu	Gln	His	Glu	Thr	Ser	Val	Ala	Arg	Tyr	Cys	Asp	Val	
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Ala	Pro	Ser	Lys	His												
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Met	Ala	Ser	Met	Ile	Lys	Asp	Val	Met	Arg	Phe	Lys	Met	Asn	Met	Glu	
1				5					10					15		
ggg	acg	gtt	aac	ggc	cac	cac	ttc	aag	tgt	gaa	gct	gat	gga	gag	ggc	96
Gly	Thr	Val	Asn	Gly	His	His	Phe	Lys	Cys	Glu	Ala	Asp	Gly	Glu	Gly	
			20					25					30			
aag	cca	tat	gaa	ggg	act	caa	ctc	gga	aga	ata	agg	gtc	acc	gaa	ggc	144

Lys	Pro	Tyr	Glu	Gly	Thr	Gln	Leu	Gly	Arg	Ile	Arg	Val	Thr	Glu	Gly	
		35					40					45				
ggg	cca	ttg	ccg	ttt	tct	ttc	gac	atc	ttg	tcg	act	tgc	act	caa	tat	192
Gly	Pro	Leu	Pro	Phe	Ser	Phe	Asp	Ile	Leu	Ser	Thr	Cys	Thr	Gln	Tyr	
	50					55					60					
gga	agc	aag	acc	atg	atc	aag	cat	ttg	tcc	ggg	att	cca	gac	tac	ttt	240
Gly	Ser	Lys	Thr	Met	Ile	Lys	His	Leu	Ser	Gly	Ile	Pro	Asp	Tyr	Phe	
65					70					75					80	
aag	cag	tct	ttt	cca	gaa	gga	ttc	acc	tgg	gaa	agg	aac	aca	atc	tat	288
Lys	G1n	Ser	Phe	Pro	Glu	G1y	Phe	Thr	Trp	Glu	Arg	Asn	Thr	Ile	Tyr	
				85					90					95		
gag	gat	ggt	ggt	cac	ctc	aca	gct	cac	caa	gac	act	agt	ctt	aag	gga	336
Glu	Asp	Gly	Gly	His	Leu	Thr	Ala	His	Gln	Asp	Thr	Ser	Leu	Lys	G1y	
			100					105					110			
gat	tgc	ctg	atc	tat	aaa	gtg	aaa	gtc	ctt	gga	ggt	aat	ttt	cct	gcc	384
Asp	Cys	Leu	Ile	Tyr	Lys	Val	Lys	Val	Leu	Gly	Gly	Asn	Phe	Pro	Ala	
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aat	ggc	cct	gtg	atg	cag	aag	aag	agc	aaa	gga	tgg	gaa	ccc	tgt	acc	432
Asn	Gly	Pro	Val	Met	Gln	Lys	Lys	Ser	Lys	Gly	Trp	Glu	Pro	Cys	Thr	
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gaa	atg	ctt	tat	cca	cgt	gat	gga	gtg	ctt	tgt	ggc	caa	aca	ttg	atg	480
Glu	Met	Leu	Tyr	Pro	Arg	Asp	Gly	Val	Leu	Cys	Gly	G1n	Thr	Leu	Met	
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gca	ctt	aaa	tgc	gcc	aat	ggt	aaa	aat	ctg	act	tgc	cag	cta	aga	tct	528
Ala	Leu	Lys	Cys	Ala	Asn	Gly	Lys	Asn	Leu	Thr	Cys	Gln	Leu	Arg	Ser	
				165					170					175		
act	tac	agg	tcc	aga	aaa	caa	gcc	agt	gca	ttg	cag	aca	cca	ggc	ttc	576

Thr Tyr Arg Ser Arg Lys Gln Ala Ser Ala Leu Gln Thr Pro Gly Phe 180 185 190 cat ttc gga gac cat cgt att gag ata ctc aag gaa gca gag ggc ggc 624 His Phe Gly Asp His Arg Ile Glu Ile Leu Lys Glu Ala Glu Gly Gly 195 200 205 aat tac ttt gag cag cac gaa aca tca gtc gcc agg tac tgt gat gtt 672 Asn Tyr Phe Glu Gln His Glu Thr Ser Val Ala Arg Tyr Cys Asp Val 210 215 220 690 gcc ccg tca aag cac tga Ala Pro Ser Lys His 225 <210> 3 <211> 21 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic DNA <400> 3 gaaggrtgyg tcaayggrca y 21 <210> 4 <211> 23 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic DNA

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<210> 5

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<223> Description of Artificial Sequence: Synthetic DNA

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19

36

<210> 7

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<212> DNA

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<400> 7

ggccacgcgt cgactagtac

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<210> 8

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<212> DNA

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<223> Description of Artificial Sequence: Synthetic DNA

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<210> 9

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<212> DNA

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<223> Description of Artificial Sequence: Synthetic DNA

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<210> 10

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<212> DNA

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<220>

<223> Description of Artificial Sequence: Synthetic DNA

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<212> DNA

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